## SEQUENCE LISTING

<110> Garber, Mitchell Ehren

```
<120> METHODS AND COMPOSITIONS FOR USE IN
  EVALUATING AND TREATING NEOPLASTIC DISEASE CONDITIONS
<130> STAN-349
<150> US2005/009926
<151> 2005-03-24
<150> 60/629,527
<151> 2004-11-18
<150> 60/558,953
<151> 2004-04-02
<160> 13
<170> FastSEQ for Windows Version 4.0
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Arg Asp Ala Arg
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Glu Leu His Leu Lys Pro His Leu Glu Gly Ala Ala Phe Arg Asp His
                                                         15
                                     10
Gln
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<400> 3
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                                                         15
                                     10
 1
Leu Asn
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Arg Glu Thr Ile Pro Ala Lys Leu Val Gln Ser Thr Leu Ser Asp Leu
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Arg
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Asp Pro Ala Lys Val Gln Ser Leu Val Asp Thr Ile Arg Glu Asp Pro
                                                          15
                                     10
Asp
<210> 6
<211> 15
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Arg Val Ala Ala Lys Arg Leu Lys Glu Gly Asp Thr Met Met Gly
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                                     10
 1
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Lys Thr Val Glu Ser Leu Glu Glu Thr Leu Lys Lys Ala Ser Pro Asp
                                     10
Gly Tyr Asp
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<211> 17
<212> PRT
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Thr Thr His Ser Ile Ser Asp Gly Lys Asp Leu Glu Lys Leu Leu Thr
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Glu
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Pro Ser Pro Tyr Glu Gly Tyr Arg
            20
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<211> 23
<212> PRT
<213> human
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Arg His Gln Leu Leu Cys Phe Lys Glu Asp Cys Gln Ala Val Phe Gln
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Asp Leu Glu Gly Val Glu Lys
            20
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<211> 19
<212> PRT
<213> human
<400> 11
Asp Leu Glu Val Lys Asp Trp Met Gln Lys Lys Arg Arg Gly Leu Arg
                 5
                                     10
                                                         15
 1
Asn Ser Arg
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<212> DNA
<213> human
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ggatccatgg aagctgcaga tgcctccagg
                                                                    30
<210> 13
<211> 71
<212> DNA
<213> human
<400> 13
accggtgtgc atcctccgcc gccgcatcct ccgccgccgg cggctggggc ttcgttggac 60
ccaatcccgt t
                                                                    71
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